

Kam

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RAW SEQUENCE LISTING

DATE: 08/15/2002

PATENT APPLICATION: US/09/595,947C

TIME: 14:27:17

Input Set : A:\Pto.raw

Output Set: N:\CRF4\08152002\I595947C.raw

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1 <110> APPLICANT: ICARD-LIEPKALNS, Christine
2     MALLET, Jacques
3     RAVASSARD, Philippe
4 <120> TITLE OF INVENTION: POLYPEPTIDES OF THE "BASIC-HELIX-LOOP-HELIX" bHLH
5     FAMILY,
6     CORRESPONDING NUCLEIC ACID SEQUENCES
7 <130> FILE REFERENCE: ST96042AUS
C--> 8 <140> CURRENT APPLICATION NUMBER: US/09/595,947C
9     <141> CURRENT FILING DATE: 2000-06-16
10    <150> PRIOR APPLICATION NUMBER: FR96/15651
11    <151> PRIOR FILING DATE: 1996-12-19
12    <150> PRIOR APPLICATION NUMBER: PCT/FR97/02368
13    <151> PRIOR FILING DATE: 1997-12-19
14    <150> PRIOR APPLICATION NUMBER: US09/331,356
15    <151> PRIOR FILING DATE: 1997-12-19
16    <160> NUMBER OF SEQ ID NOS: 28
17    <170> SOFTWARE: PatentIn version 3.0
19    <210> SEQ ID NO: 1
20    <211> LENGTH: 1460
21    <212> TYPE: DNA
22    <213> ORGANISM: Rattus norvegicus
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25      gcagcccggc aggcacgctc ctggtccggg cagagcagat aaagcgtgcc aggggacaca      120
26      cgattagcag ctcaagaagtc cctctgggtc tcaccactgc acagaggccg aggaccccct      180
27      ccgagcttct ttgctgcctc cagacgcaat ttactccagg cgagggcgcc tgcagctcag      240
28      caaaacttcg aagcgagcag aggggttcag ctatccaccg ctgcttgact ctgaccaccc      300
29      gcagctctct gttcttttga gcccgagta actaggtaac atttaggaac ctccaaaggg      360
30      tagaagaggg gagtgggtgg gcgtactcta gtcccgcgtg gagtgcctc taagtcagag      420
31      actgtcacac cccccttcca ttttttccca acctcaggat ggcgcctcat cccttggatg      480
32      cgcccaccat ccaagtgtcc caagagaccc agcaaccctt tcccggagcc tccgaccacg      540
33      aagtgtcag ttccaattcc accccaccta gcccactct cgtaccgagg gactgctccg      600
34      aagcagaagc aggtgactgc cgagggacat cgaggaagct ccgtgcgcgg cgcggagggc      660
35      gcaacaggcc caagagcgag ttggcactga gcaagcagcg acgaagccgg cgcaagaagg      720
36      ccaacgaccg ggagcgcaac cgcattgcaca accttaactc cgcgctggat gcgctgcgcg      780
37      gtgtcctgcc caccttcccg gatgacgcca aacttacaaa gatcgagacc ctgcgcttcg      840
38      cccacaacta catttgggca ctgactcaga cgctgcgcac agcggaccac agcttctacg      900
39      gccccgagcc ccctgtgccc tgtggggagc tgggaagccc gggagggggc tccagcggcg      960
40      actggggctc tatctactcc ccagtttccc aagctggtag cctgagcccc acagcctcat      1020
41      tggaggagtt ccctggcctg caggtgcccc gctccccatc ctgtctgctc ccgggcaccc      1080
42      tgggtgttctc agacttcttg tgaagggccc aaacaggccc tgggcgggtgg gcgctggcag      1140
43      aaagggaggg agtcagagct gtctgaaatg gaaggtagtg gaggcactcg agcatctcgc      1200
44      cccttctggc tticattagt caggtccctg atttaaccag gattcgacaca gttccttgc      1260

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45      gctgtgcgtg cacaaaggac attgcaggct gatctcctct taaccctcct cagtgtggcc 1320
46      acctcaaaact ccgcgtccaa gcagaggaga gccgtagcac taaatagttg ggagactccc 1380
47      atacttcctg gtgactccgc cctctttcaa atctgcgggc ctccaaccac cgctttctcc 1440
48      agagtgcacct aatccagtgt                                     1460
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51 <211> LENGTH: 24
52 <212> TYPE: PRT
53 <213> ORGANISM: Artificial Sequence
54 <220> FEATURE:
55 <223> OTHER INFORMATION: peptide fragment of bHLH protein
56 <400> SEQUENCE: 2
57      Ala Ala Thr Lys His Gly Met Gly Ile Gly Ala Gly Cys Gly Cys Ile
58      1              5              10              15
59      Asp Lys Cys Gly Cys Arg Tyr Gly
60              20
62 <210> SEQ ID NO: 3
63 <211> LENGTH: 24
64 <212> TYPE: PRT
65 <213> ORGANISM: Artificial Sequence
66 <220> FEATURE:
67 <223> OTHER INFORMATION: peptide fragment of bHLH protein
68 <400> SEQUENCE: 3
69      Gly Gly Cys Ser Arg Asp Thr Tyr Thr Cys Ala Gly Gly Gly Thr Ser
70      1              5              10              15
71      Tyr Asx Gly Ala Tyr Cys Thr Thr
72              20
74 <210> SEQ ID NO: 4
75 <211> LENGTH: 25
76 <212> TYPE: DNA
77 <213> ORGANISM: Artificial Sequence
78 <220> FEATURE:
79 <223> OTHER INFORMATION: primer
80 <400> SEQUENCE: 4
81      aaccttaact ccgcgctgga tgcgc                                     25
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84 <211> LENGTH: 18
85 <212> TYPE: DNA
86 <213> ORGANISM: Artificial Sequence
87 <220> FEATURE:
88 <223> OTHER INFORMATION: primer
89 <400> SEQUENCE: 5
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93 <211> LENGTH: 6
94 <212> TYPE: DNA
95 <213> ORGANISM: Artificial Sequence
96 <220> FEATURE:
97 <223> OTHER INFORMATION: DNA sequence of E box
98 <400> SEQUENCE: 6

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102 <211> LENGTH: 6
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104 <213> ORGANISM: Artificial Sequence
105 <220> FEATURE:
106 <223> OTHER INFORMATION: DNA sequence of mutated E box
107 <400> SEQUENCE: 7
108      tccgtg                                     6
110 <210> SEQ ID NO: 8
111 <211> LENGTH: 214
112 <212> TYPE: PRT
113 <213> ORGANISM: Rattus norvegicus
114 <400> SEQUENCE: 8
115      Met Ala Pro His Pro Leu Asp Ala Pro Thr Ile Gln Val Ser Gln Glu
116      1              5              10              15
117      Thr Gln Gln Pro Phe Pro Gly Ala Ser Asp His Glu Val Leu Ser Ser
118      20              25              30
119      Asn Ser Thr Pro Pro Ser Pro Thr Leu Val Pro Arg Asp Cys Ser Glu
120      35              40              45
121      Ala Glu Ala Gly Asp Cys Arg Gly Thr Ser Arg Lys Leu Arg Ala Arg
122      50              55              60
123      Arg Gly Gly Arg Asn Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
124      65              70              75              80
125      Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
126      85              90              95
127      His Asn Leu Asn Ser Ala Leu Asp Ala Leu Arg Gly Val Leu Pro Thr
128      100             105             110
129      Phe Pro Asp Asp Ala Lys Leu Thr Lys Ile Glu Thr Leu Arg Phe Ala
130      115             120             125
131      His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
132      130             135             140
133      Ser Phe Tyr Gly Pro Glu Pro Pro Val Pro Cys Gly Glu Leu Gly Ser
134      145             150             155             160
135      Pro Gly Gly Gly Ser Ser Gly Asp Trp Gly Ser Ile Tyr Ser Pro Val
136      165             170             175
137      Ser Gln Ala Gly Ser Leu Ser Pro Thr Ala Ser Leu Glu Glu Phe Pro
138      180             185             190
139      Gly Leu Gln Val Pro Ser Ser Pro Ser Cys Leu Leu Pro Gly Thr Leu
140      195             200             205
141      Val Phe Ser Asp Phe Leu
142      210
144 <210> SEQ ID NO: 9
145 <211> LENGTH: 1330
146 <212> TYPE: DNA
147 <213> ORGANISM: Homo sapiens
148 <400> SEQUENCE: 9
149      cctcggaccc cattctctct tcttttctcc tttggggctg gggcaactcc caggcggggg      60
150      cgctgcagc tcagctgaac ttggcgacca gaagcccgt gagctcccca cggccctcgc      120

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151      tgctcatcgc tctctattct tttgcgccgg tagaaaggta atatttgagg gccttcgagg      180
152      gacggggcagg ggaaaaggagg atcctctgac ccagcggggg ctggggaggat ggctgttttt      240
153      gttttttccc acctagcctc ggaatcgccg actgcgccgt gacgggactca aactttacct      300
154      tccctctgac cccgccgtag gatgacgcct caaccctcgg gtgcgcccac tgtccaagtg      360
155      acccgtgaga cggagcggtc cttccccaga gcctcggaag acgaagtgac ctgccccacg      420
156      tccgccccgc ccagccccac tcgcacaccg gggaactgcg cagaggcgga agagggaggc      480
157      tgccgagggg ccccgaggaa gctccgggca cggcgcgggg gacgcagccg gcctaagagc      540
158      gagttggcac tgagcaagca gcgacggagt cggcgaaaaga aggccaacga ccgcgagcgc      600
159      aatcgaatgc acgacctcaa ctcggcactg gacgcccctgc gcggtgtcct gcccaccttc      660
160      ccagacgacg cgaagctcac caagatcgag acgctgcgct tcgcccacaa ctacatctgg      720
161      gcgctgactc aaacgctgcg catagcggac cacagcttgt acgcgctgga gccgcggcg      780
162      ccgcaactgc gggagctggg cagcccaggc ggtccccccg gggactgggg gtccctctac      840
163      tccccagtct cccaggctgg cagcctgagt cccgcgcgct cgctggagga gcgacccggg      900
164      ctgctggggg ccacctcttc cgctgtcttg agcccaggca gtctggcttt ctcagatttt      960
165      ctgtgaaaag acctgtctgt cgctgggctg tgggtgctaa gggtaaggga gagggaggga      1020
166      gccgggagcc gtagagggtg gccgacggcg gcggccctca aaagcacttg ttcttctgc      1080
167      ttctccctag ctgacccctg gccggcccag gcctccacgg gggcggtagg ctgggttcat      1140
168      tccccggccc tccgagccgc gccaacgcac gcaacccttg ctgctgcccg cgcgaagtgg      1200
169      gcattgcaaa gtgcgctcat tttaggcctc ctctctgcca ccacccata atcccattca      1260
170      aagaatacta gaatggtagc actaccggc cggagccgcc caccgtcttg ggtcgcccta      1320
171      ccctcactca                                     1330

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173 <210> SEQ ID NO: 10

174 <211> LENGTH: 214

175 <212> TYPE: PRT

176 <213> ORGANISM: Homo sapiens

177 <400> SEQUENCE: 10

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178      Met Thr Pro Gln Pro Ser Gly Ala Pro Thr Val Gln Val Thr Arg Glu
179      1          5          10          15
180      Thr Glu Arg Ser Phe Pro Arg Ala Ser Glu Asp Glu Val Thr Cys Pro
181      20          25          30
182      Thr Ser Ala Pro Pro Ser Pro Thr Arg Thr Pro Gly Asn Cys Ala Glu
183      35          40          45
184      Ala Glu Glu Gly Gly Cys Arg Gly Ala Pro Arg Lys Leu Arg Ala Arg
185      50          55          60
186      Arg Gly Gly Arg Ser Arg Pro Lys Ser Glu Leu Ala Leu Ser Lys Gln
187      65          70          75          80
188      Arg Arg Ser Arg Arg Lys Lys Ala Asn Asp Arg Glu Arg Asn Arg Met
189      85          90          95
190      His Asp Leu Asn Ser Ala Leu Asp Ala Leu Arg Gly Val Leu Pro Thr
191      100         105         110
192      Phe Pro Asp Asp Ala Lys Leu Thr Lys Ile Glu Thr Leu Arg Phe Ala
193      115         120         125
194      His Asn Tyr Ile Trp Ala Leu Thr Gln Thr Leu Arg Ile Ala Asp His
195      130         135         140
196      Ser Leu Tyr Ala Leu Glu Pro Pro Ala Pro His Cys Gly Glu Leu Gly
197      145         150         155         160
198      Ser Pro Gly Gly Pro Pro Gly Asp Trp Gly Ser Leu Tyr Ser Pro Val
199      165         170         175
200      Ser Gln Ala Gly Ser Leu Ser Pro Ala Ala Ser Leu Glu Glu Arg Pro

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201          180          185          190
202      Gly Leu Leu Gly Ala Thr Ser Ser Ala Cys Leu Ser Pro Gly Ser Leu
203          195          200          205
204      Ala Phe Ser Asp Phe Leu
205          210
207 <210> SEQ ID NO: 11
208 <211> LENGTH: 18
209 <212> TYPE: DNA
210 <213> ORGANISM: Artificial Sequence
211 <220> FEATURE:
212 <223> OTHER INFORMATION: primer
213 <400> SEQUENCE: 11
214      caacgaccgg cagcgcaa
216 <210> SEQ ID NO: 12
217 <211> LENGTH: 24
218 <212> TYPE: DNA
219 <213> ORGANISM: Artificial Sequence
220 <220> FEATURE:
221 <223> OTHER INFORMATION: primer
222 <400> SEQUENCE: 12
223      gcccagatgt agttgtgggc gaag
225 <210> SEQ ID NO: 13
226 <211> LENGTH: 60
227 <212> TYPE: DNA
228 <213> ORGANISM: Artificial Sequence
229 <220> FEATURE:
230 <223> OTHER INFORMATION: primer
231 <220> FEATURE:
232 <221> NAME/KEY: misc_feature
233 <223> OTHER INFORMATION: n=a or t or g or c
234 <400> SEQUENCE: 13
W--> 235      atcggttgaga ctcgtaccag cagagtcacg agagagacta cacggtactg gnnnnnnnnnn
237 <210> SEQ ID NO: 14
238 <211> LENGTH: 20
239 <212> TYPE: DNA
240 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: primer
243 <400> SEQUENCE: 14
244      agacgacgcg aagctcacca
246 <210> SEQ ID NO: 15
247 <211> LENGTH: 24
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: primer
252 <400> SEQUENCE: 15
253      gctcaccaag atcgagacgc tgcg
255 <210> SEQ ID NO: 16

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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 08/15/2002
PATENT APPLICATION: US/09/595,947C TIME: 14:27:18

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:13; N Pos. 52,53,54,55,56,57,58,59,60